

Exhibit 1

Blast Result

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 NCBI Blast 2 Sequences results

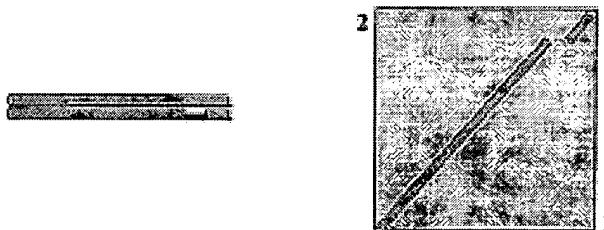
Published Entrez BLAST OMIM Taxonomy Structure

BLAST 2 SEQUENCES RESULTS VERSION BLASTP 2.2.17 [Aug-19-2007]

Matrix BLOSUM62 gap open: 11 gap extension: 1
X-dropoff: 0 expect: 10.000 wordsize: 3 Filter View option Standard
Masking character option X for protein, n for nucleotide Masking color option Black
 Show CDS translation

Sequence 1: unnamed protein product
Length = 71 (1 .. 71)

Sequence 2: unnamed protein product
Length = 64 (1 .. 64)



NOTE: Bitscore and expect value are calculated based on the size of the nr database.

Score = 94.0 bits (232), Expect = 2e-18
Identities = 60/70 (85%), Positives = 61/70 (87%), Gaps = 2/70 (10%)

Query	1	MVLRQLSRKASVVKVQTKTQGTTKARAGCILILFLLDFLLNPFCTGEGEDSVDGEKRAQRESCLTQ	60
		MVLRQLSRKASVVKVQTKTQGTTKARAGCILILFLLDFLLNPFCTGEGEDSVDGEKRAQRESCLTQ	
Subject	1	MVLRQLSRKASVVKVQTKTQGTTKARAGCILILFLLDFLLNPFCTGEGEDSVDGEKRAQRESCLTQ	56
		MVLRQLSRKASVVKVQTKTQGTTKARAGCILILFLLDFLLNPFCTGEGEDSVDGEKRAQRESCLTQ	
Query	61	TYBALPEPKA 70 SEQ. IDN: /	
		ALPEPKA	
Subject	57	---ALPEPKA 63 SEQ. IDN: /	

CPU time: 0.03 user secs., 0.01 sys. secs. 0.04 total secs.